

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/531,870
Source: PT/10
Date Processed by STIC: 9/12/05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebs/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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Revised 01/24/05



PCT

RAW SEQUENCE LISTING

DATE: 09/12/2005

PATENT APPLICATION: US/10/531,870

TIME: 13:21:40

Input Set : A:\764765_1.txt

Output Set: N:\CRF4\09122005\J531870.raw

3 <110> APPLICANT: FIT Biotech Oyj Plc.
 5 <120> TITLE OF INVENTION: Novel selection system
 7 <130> FILE REFERENCE: 2031002
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/531,870
 C--> 9 <141> CURRENT FILING DATE: 2005-04-19
 9 <160> NUMBER OF SEQ ID NOS: 29
 11 <170> SOFTWARE: PatentIn version 3.1
 13 <210> SEQ ID NO: 1
 14 <211> LENGTH: 780
 15 <212> TYPE: DNA
 16 <213> ORGANISM: Escherichia coli
 18 <400> SEQUENCE: 1
 19 gtttcgtttg attggctgtg gttttataca gtcattactg cccgtaatat gccttcgcgc 60
 21 catgcttacg cagatagtgt ttatccagca gcgtttgctg catatccggt aactgcggcg 120
 23 ctaactgacg gcagaatatc cccatataag cgacctcttc cagcacgatg gcgttatgca 180
 25 ccgcaccttc ggcatttttg ccccatgcaa acgggcccgtg ggaatggacc agaacgccgg 240
 27 gcatttgccg tgcacgata ccctgttttt caaagggttc tacgatgacg ttaccgggtt 300
 29 cccactcata ttcgccgttg atttctgcgt cggtcatttt gcgggtgcag ggaatgggtg 360
 31 cgtagaaata gtcggcgtgg gtggtgccgg ttgctggaat cgactgaccc gcctgcgccc 420
 33 agatgggtggc gtggcgcgag tgcgtatgca caatgccgcc aatggagggg aatgcctgat 480
 35 agagcagccg gtgagttggc gtgtcggagg agggcttttt cgtaccttca accacttcac 540
 37 cggtttcgat gctaaccacg accatatcgt cagcggtcac gacgctgtaa tcgacgccgg 600
 39 aaggtttgat cacaaagacg ccgcgctcgc gatcaacggc gctgacgttg ccccatgtga 660
 41 gcgtgaccag gttgtgtttt ggcagcgcca ggttggcttc taatacctgg cgtttgagat 720
 43 cttctaacat gttgactcct tcgtgccgga tgcgctttgc ttatccggcc tacaaaaatcg 780
 46 <210> SEQ ID NO: 2
 47 <211> LENGTH: 76
 48 <212> TYPE: DNA
 49 <213> ORGANISM: Artificial Sequence
 51 <220> FEATURE:
 52 <223> OTHER INFORMATION: Primer
 54 <400> SEQUENCE: 2
 55 cgccatgggt ctcattgttg acagcttatc atcgataagc tttaatgcgg tagtttagca 60
 57 cgaaggagtc aacatg 76
 60 <210> SEQ ID NO: 3
 61 <211> LENGTH: 64
 62 <212> TYPE: DNA
 63 <213> ORGANISM: Artificial Sequence
 65 <220> FEATURE:
 66 <223> OTHER INFORMATION: Primer
 68 <400> SEQUENCE: 3
 69 cgccatggac tagtaaaaaa aagcccgcgc attaggcggg ctgtcattac tgcccgtaat 60
 71 atgc 64

Does Not Comply
Corrected Diskette Needed

pg 4-6

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TIME: 13:21:40

Input Set : A:\764765_1.txt

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74 <210> SEQ ID NO: 4
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80 <223> OTHER INFORMATION: Primer
82 <400> SEQUENCE: 4
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85 tagcacgaag gagtcaacat g                                     81
88 <210> SEQ ID NO: 5
89 <211> LENGTH: 58
90 <212> TYPE: DNA
91 <213> ORGANISM: Artificial Sequence
93 <220> FEATURE:
94 <223> OTHER INFORMATION: Primer
96 <400> SEQUENCE: 5
97 cgccatggaa aaaaaagccc gctcattagg cgggctgtca ttactgcccg taatatgc      58
100 <210> SEQ ID NO: 6
101 <211> LENGTH: 22
102 <212> TYPE: DNA
103 <213> ORGANISM: Artificial Sequence
105 <220> FEATURE:
106 <223> OTHER INFORMATION: Primer
108 <400> SEQUENCE: 6
109 gccagggttt tcccagtcac ga                                     22
112 <210> SEQ ID NO: 7
113 <211> LENGTH: 24
114 <212> TYPE: DNA
115 <213> ORGANISM: Artificial Sequence
117 <220> FEATURE:
118 <223> OTHER INFORMATION: Primer
120 <400> SEQUENCE: 7
121 gagcggataa caatttcaca cagg                                     24
124 <210> SEQ ID NO: 8
125 <211> LENGTH: 22
126 <212> TYPE: DNA
127 <213> ORGANISM: Artificial Sequence
129 <220> FEATURE:
130 <223> OTHER INFORMATION: Primer
132 <400> SEQUENCE: 8
133 ccaactcacc ggctgctcta tc                                     22
136 <210> SEQ ID NO: 9
137 <211> LENGTH: 24
138 <212> TYPE: DNA
139 <213> ORGANISM: Artificial Sequence
141 <220> FEATURE:
142 <223> OTHER INFORMATION: Primer
144 <400> SEQUENCE: 9
145 aatgccgaag atgcggtgca taac                                     24

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Input Set : A:\764765_1.txt

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148 <210> SEQ ID NO: 10
149 <211> LENGTH: 20
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151 <213> ORGANISM: Artificial Sequence
153 <220> FEATURE:
154 <223> OTHER INFORMATION: Primer
156 <400> SEQUENCE: 10
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160 <210> SEQ ID NO: 11
161 <211> LENGTH: 20
162 <212> TYPE: DNA
163 <213> ORGANISM: Artificial Sequence
165 <220> FEATURE:
166 <223> OTHER INFORMATION: Primer
168 <400> SEQUENCE: 11
169 ggttgctgga atcgactgac 20
172 <210> SEQ ID NO: 12
173 <211> LENGTH: 66
174 <212> TYPE: DNA
175 <213> ORGANISM: Artificial Sequence
177 <220> FEATURE:
178 <223> OTHER INFORMATION: Primer
180 <400> SEQUENCE: 12
181 ctcaaagcc caggtattag aagccaacct ggcgctgcc aaacacgtgt aggctggagc 60
183 tgcttc 66
186 <210> SEQ ID NO: 13
187 <211> LENGTH: 60
188 <212> TYPE: DNA
189 <213> ORGANISM: Artificial Sequence
191 <220> FEATURE:
192 <223> OTHER INFORMATION: Primer
194 <400> SEQUENCE: 13
195 ggtttgatca caaagacgcc gcgctcgca tcaacggcgc attccgggga tccgtcgacc 60
198 <210> SEQ ID NO: 14
199 <211> LENGTH: 20
200 <212> TYPE: DNA
201 <213> ORGANISM: Artificial Sequence
203 <220> FEATURE:
204 <223> OTHER INFORMATION: Primer
206 <400> SEQUENCE: 14
207 cggcacgaag gagtcaacat 20
210 <210> SEQ ID NO: 15
211 <211> LENGTH: 20
212 <212> TYPE: DNA
213 <213> ORGANISM: Artificial Sequence
215 <220> FEATURE:
216 <223> OTHER INFORMATION: Primer
218 <400> SEQUENCE: 15
219 tgatagagca gccggtgagt 20

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Input Set : A:\764765_1.txt

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222 <210> SEQ ID NO: 16
223 <211> LENGTH: 20
224 <212> TYPE: DNA
225 <213> ORGANISM: Artificial Sequence
227 <220> FEATURE:
228 <223> OTHER INFORMATION: Primer
230 <400> SEQUENCE: 16
231 tcagatcctt ggcggcaaga                20
234 <210> SEQ ID NO: 17
235 <211> LENGTH: 20
236 <212> TYPE: DNA
237 <213> ORGANISM: Artificial Sequence
239 <220> FEATURE:
240 <223> OTHER INFORMATION: Primer
242 <400> SEQUENCE: 17
243 tgtaatcgac gccggaaggt                20
246 <210> SEQ ID NO: 18
247 <211> LENGTH: 1030
248 <212> TYPE: DNA
249 <213> ORGANISM: Artificial
W--> 251 <220> FEATURE:
W--> 251 <223> OTHER INFORMATION:
W--> 251 <400> 18
252 ggatccgacc ggcaacggta cagatccgac cggcaacggt acagatccga ccggcaacgg      60
254 tcagatccga ccggcaacgg tacagatccg accggcaacg gtacagatcc gaccggcaac      120
256 ggtacagatc cgaccggcaa cggtagatcc ccgaccggca acggtacaga tccgaccggc      180
258 aacgggtacag atccgaccgg caacgggtaca gatcccccta gcgaattgac tagttctcat      240
260 gtttgacagc ttatcatcga taagctttaa tgcggtagtt tagcacgaag gagtcaacat      300
262 gttagaagat ctcaaacgcc aggtattaga agccaacctg gcgctgccaa aacacaacct      360
264 ggtcacgctc acatggggca acgtcagcgc cgttgatcgc gagcgcgcg tctttgtgat      420
266 caaaccttcc ggcgtcgatt acagcgtcat gaccgctgac gatatggctg tggtagcat      480
268 cgaacccggt gaagtgggtg aaggtacgaa aaagccctcc tccgacacgc caactcaccg      540
270 gctgctctat caggcattcc cctccattgg cggcattgtg catacgcact cgcgccacgc      600
272 caccatctgg gcgcaggcgg gtcagtcgat tccagcaacc ggcaccaccc acgccgacta      660
274 tttctacggc accattccct gcacccgcaa aatgaccgac gcagaaatca acggcgaata      720
276 tgagtgggaa accggtaacg tcatcgtaga aacctttgaa aaacagggta tcgatgcagc      780
278 gcaaagtccc ggcgttctgg tccattccca cgccccgttt gcatggggca aaaatgccga      840
280 agatgcggtg cataacgcca tcgtgctgga agaggtcgct tatatgggga tattctgccc      900
282 tcagttagcg ccgcagttac cggatatgca gcaaacgctg ctggataaac actatctgcg      960
284 taagcatggc gcgaaggcat attacgggca gtaatgacag cccgcctaata gagcgggctt     1020
286 ttttttccat                                1030
289 <210> SEQ ID NO: 19
290 <211> LENGTH: 696
291 <212> TYPE: DNA
292 <213> ORGANISM: Escherichia coli
294 <400> SEQUENCE: 19
295 atgttagaag atctcaaacg ccagggtatta gaagccaacc tggcgctgcc aaaacacaac      60
297 ctggtcacgc tcacatgggg caacgtcagc gccgttgatc gcgagcgcg cgtctttgtg      120
299 atcaaacctt ccggcgctga ttacagcgtc atgaccgctg acgatatggt cgtggttagc      180

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needs explanation (see p.6 for error explanation)

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301 atcgaaaccg gtgaagtggg tgaaggtacg aaaaagccct cctccgacac gccaaactcac 240
303 cggtctgtct atcaggcatt cccctccatt ggcggcattg tgcatacgca ctccgcccac 300
305 gccaccatct gggcgaggc ggggtcagtc attccagcaa ccggcaccac ccacgccgac 360
307 tttttctacg gcaccattcc ctgcacccgc aaaatgaccg acgcagaaat caacggcgaa 420
309 tatgagtggg aaaccggtaa cgtcatcgta gaaacctttg aaaaacaggg tatcgatgca 480
311 gcgcaaagtc ccggcggttct ggtccattcc cacggcccgt ttgcatgggg caaaaatgcc 540
313 gaagatgcgg tgcataacgc catcgtgctg gaagaggtcg cttatatggg gatattctgc 600
315 cgtcagttag cgccgcagtt accggatatg cagcaaacgc tgctggataa aactatctgc 660
317 cgtaagcatg gcgcgaaggc atattacggg cagtaa 696

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320 <210> SEQ ID NO: 20

321 <211> LENGTH: 687

322 <212> TYPE: DNA

323 <213> ORGANISM: Escherichia coli

325 <400> SEQUENCE: 20

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326 atgcaaaagc taaaacagca ggtattttaa gccaacatgg agctgccgcg ctacgggctg 60
328 gtgaccttta cctggggcaa cgtcagcgct atcgaccgcg aacgcgggct ggtggtgatc 120
330 aagcccagcg gcgttgcccta cgaaaccatg aaagcggccg atatggtggt ggttgatatg 180
332 agcggcaagg tgggtggaagg ggagtatcgc ccactctccg acactgcgac gcactctgaa 240
334 ctctaccgtc gttacccgtc gcttggtggc attgtccata cccactccac tcatgccacc 300
336 gcatggggcg aggcgggggct ggcgatcccc gcgttaggca ccacgcacgc cgactacttc 360
338 tttggcgaca ttccgtgtac gcgcggggtta agcgaagaag aggtgcaggg cgagtatgaa 420
340 ctgaacaccg gcaaagtgat tatcgaaacg ctgggcaacg ccgagccgct gcatacgccg 480
342 ggaattgtgg tgtatcagca cgggccgttc gcctggggga aagatgctca cgatgcggtg 540
344 cataacgcgg tggatgagga agaagtggcg aaaatggcgt ggattgcccg cggcattaac 600
346 ccacaactca atcacatcga cagcttcctg atgaataaac acttcatgcg taaacacggt 660
348 cctaacgctt attacgggca gaagtag 687

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351 <210> SEQ ID NO: 21

352 <211> LENGTH: 65

353 <212> TYPE: DNA

354 <213> ORGANISM: Artificial

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W--> 356 <223> OTHER INFORMATION:

W--> 356 <400> 21

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357 cagcaggtat ttgaagccaa catggagctg ccgcgctacg ggctggtgta ggctggagct 60
359 gcttc 65

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362 <210> SEQ ID NO: 22

363 <211> LENGTH: 66

364 <212> TYPE: DNA

365 <213> ORGANISM: Artificial

W--> 367 <220> FEATURE:

W--> 367 <223> OTHER INFORMATION:

W--> 367 <400> 22

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368 aaacggctgc ggaattagac cagttatctc ccgaggaagg aaattaattc cggggatccc 60
370 tcgacc 66

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373 <210> SEQ ID NO: 23

374 <211> LENGTH: 21

375 <212> TYPE: DNA

376 <213> ORGANISM: Artificial

W--> 378 <220> FEATURE:

*error occurs in subsequent sequences
(see p.6)*

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/531,870

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Input Set : A:\764765_1.txt
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:18,21,22,23,24,26,27,28,29

Use of <220> Feature(NEW RULES):

error explanation
Sequence(s) are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence"
or "Unknown". Please explain source of genetic material in <220> to <223>
section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32)
(Sec.1.823 of new Rules)

Seq#:18,21,22,23,24,26,27,28,29

VERIFICATION SUMMARY

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Input Set : A:\764765_1.txt

Output Set: N:\CRF4\09122005\J531870.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:251 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:18, <213>
ORGANISM:Artificial
L:251 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:18, <213>
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L:251 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:18,Line#:251
L:356 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:21, <213>
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L:367 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:22,Line#:367
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L:427 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:26, <213>
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